## SEQUENCE LISTING

```
<110> Banerjee, Subhashis
      Taylor, Lori K
      Spiegler, Clive E
      Tracey, Daniel E
      Chartash, Elliot K
      Hoffman, Rebecca S
      Barchuk, William T
      Yan, Philip
      Murtaza, Anwar
Salfeld, Jochen G
      Fischkoff, Steven
<120> TREATMENT OF PULMONARY DISORDERS USING TNF\alpha INHIBITORS
<130> BPI-189
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<141>
<150> 60/397,275
<151> 2002-07-19
<150> 60/411,081
<151> 2002-09-16
<150> 60/417,490
<151> 2002-10-10
<150> 60/455,777
<151> 2003-03-18
<160> 37
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Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly
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                 5
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Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Gly Ile Arg Asn Tyr
                                 25
Leu Ala Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile
                             40
Tyr Ala Ala Ser Thr Leu Gln Ser Gly Val Pro Ser Arg Phe Ser Gly
                         55
Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro
                                         75
Glu Asp Val Ala Thr Tyr Tyr Cys Gln Arg Tyr Asn Arg Ala Pro Tyr
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85
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Thr Phe Gly Gln Gly Thr Lys Val Glu Ile Lys
            100
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<400> 2
Glu Val Gln Leu Val Glu Ser Gly Gly Leu Val Gln Pro Gly Arg
                5
                                    10
Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Asp Asp Tyr
            20
                                25
Ala Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
Ser Ala Ile Thr Trp Asn Ser Gly His Ile Asp Tyr Ala Asp Ser Val
Glu Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Asn Ser Leu Tyr
Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
               85
                                   90
Ala Lys Val Ser Tyr Leu Ser Thr Ala Ser Ser Leu Asp Tyr Trp Gly
                            105
Gln Gly Thr Leu Val Thr Val Ser Ser
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<210> 3
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<223> Xaa = Thr or Ala
<223> Mutated human antibody
<400> 3
Gln Arg Tyr Asn Arg Ala Pro Tyr Xaa
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<210> 4
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<221> VARIANT
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<223> Xaa = Tyr or Asn
<223> Mutated human antibody
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Val Ser Tyr Leu Ser Thr Ala Ser Ser Leu Asp Xaa
                 5
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<400> 5
Ala Ala Ser Thr Leu Gln Ser
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Ala Ile Thr Trp Asn Ser Gly His Ile Asp Tyr Ala Asp Ser Val Glu
1
                                     10
Gly
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Arg Ala Ser Gln Gly Ile Arg Asn Tyr Leu Ala
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Asp Tyr Ala Met His
1
                 5
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<400> 9
Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Ile Gly
                                    10
Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Gly Ile Arg Asn Tyr
                                25
Leu Ala Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile
Tyr Ala Ala Ser Thr Leu Gln Ser Gly Val Pro Ser Arg Phe Ser Gly
Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro
                                        75
Glu Asp Val Ala Thr Tyr Tyr Cys Gln Lys Tyr Asn Ser Ala Pro Tyr
                                   90
                85
Ala Phe Gly Gln Gly Thr Lys Val Glu Ile Lys
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<400> 10
Gln Val Gln Leu Val Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Arg
Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Asp Asp Tyr
                                25
Ala Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Asp Trp Val
                           40
Ser Ala Ile Thr Trp Asn Ser Gly His Ile Asp Tyr Ala Asp Ser Val
                       55
Glu Gly Arg Phe Ala Val Ser Arg Asp Asn Ala Lys Asn Ala Leu Tyr
                   70
                                        75
Leu Gln Met Asn Ser Leu Arg Pro Glu Asp Thr Ala Val Tyr Tyr Cys
               85
                                   90
Thr Lys Ala Ser Tyr Leu Ser Thr Ser Ser Ser Leu Asp Asn Trp Gly
Gln Gly Thr Leu Val Thr Val Ser Ser
<210> 11
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<213> Artificial Sequence
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<400> 11
Gln Lys Tyr Asn Ser Ala Pro Tyr Ala
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<210> 12
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<212> PRT
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Gln Lys Tyr Asn Arg Ala Pro Tyr Ala
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<400> 13
Gln Lys Tyr Gln Arg Ala Pro Tyr Thr
<210> 14
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Gln Lys Tyr Ser Ser Ala Pro Tyr Thr
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<400> 15
Gln Lys Tyr Asn Ser Ala Pro Tyr Thr
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Gln Lys Tyr Asn Arg Ala Pro Tyr Thr
<210> 17
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<212> PRT
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<400> 17
Gln Lys Tyr Asn Ser Ala Pro Tyr Tyr
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<400> 18
Gln Lys Tyr Asn Ser Ala Pro Tyr Asn
                 5
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Gln Lys Tyr Thr Ser Ala Pro Tyr Thr
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Gln Lys Tyr Asn Arg Ala Pro Tyr Asn
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<220>
<223> Mutated human antibody
Gln Lys Tyr Asn Ser Ala Ala Tyr Ser
                 5
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<223> Mutated human antibody
<400> 22
Gln Gln Tyr Asn Ser Ala Pro Asp Thr
                 5
<210> 23
<211> 9
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<223> Mutated human antibody
Gln Lys Tyr Asn Ser Asp Pro Tyr Thr
<210> 24
<211> 9
<212> PRT
<213> Artificial Sequence
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<223> Mutated human antibody
Gln Lys Tyr Ile Ser Ala Pro Tyr Thr
<210> 25
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<223> Mutated human antibody
<400> 25
Gln Lys Tyr Asn Arg Pro Pro Tyr Thr
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<210> 26
<211> 9
<212> PRT
<213> Artificial Sequence
<220>
<223> Mutated human antibody
<400> 26
Gln Arg Tyr Asn Arg Ala Pro Tyr Ala
<210> 27
<211> 12
<212> PRT
<213> Artificial Sequence
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<223> Mutated human antibody
<400> 27
Ala Ser Tyr Leu Ser Thr Ser Ser Ser Leu Asp Asn
            5
<210> 28
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<400> 28
Ala Ser Tyr Leu Ser Thr Ser Ser Ser Leu Asp Lys
                5
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 <400> 29
 Ala Ser Tyr Leu Ser Thr Ser Ser Ser Leu Asp Tyr
                 5
 <210> 30
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 <212> PRT
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<223> Mutated human antibody
<400> 30
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                5
<210> 31
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<223> Mutated human antibody
<400> 31
Ala Ser Tyr Leu Ser Thr Ser Phe Ser Leu Asp Tyr
                5
<210> 32
<211> 12
<212> PRT
<213> Artificial Sequence
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<223> Mutated human antibody
Ala Ser Tyr Leu Ser Thr Ser Ser Ser Leu His Tyr
                5
<210> 33
<211> 12
<212> PRT
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<400> 33
Ala Ser Phe Leu Ser Thr Ser Ser Ser Leu Glu Tyr
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<210> 34
<211> 12
<212> PRT
<213> Artificial Sequence
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<400> 34
Ala Ser Tyr Leu Ser Thr Ala Ser Ser Leu Glu Tyr
<210> 35
<211> 12
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<213> Artificial Sequence
<223> Mutated human antibody
<400> 35
Val Ser Tyr Leu Ser Thr Ala Ser Ser Leu Asp Asn
<210> 36
<211> 321
<212> DNA
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<223> Mutated human antibody
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atcacttgtc gggcaagtca gggcatcaga aattacttag cctggtatca gcaaaaacca 120
gggaaagccc ctaagctcct gatctatgct gcatccactt tgcaatcagg ggtcccatct 180
cggttcagtg gcagtggatc tgggacagat ttcactctca ccatcagcag cctacagcct 240
gaagatgttg caacttatta ctgtcaaagg tataaccgtg caccgtatac ttttggccag 300
gggaccaagg tggaaatcaa a
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<210> 37
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teetgtgegg eetetggatt eacetttgat gattatgeea tgeactgggt eeggeaaget 120
ccagggaagg gcctggaatg ggtctcagct atcacttgga atagtggtca catagactat 180
gcggactctg tggagggccg attcaccatc tccagagaca acgccaagaa ctccctgtat 240
ctgcaaatga acagtctgag agctgaggat acggccgtat attactgtgc gaaagtctcg 300
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agt
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